

# EBI plans

- Imputation with genotypes instead of haplotypes (Natalja, in progress)
- microRNAs – analysis of results from Kraken pipeline (Natalja, Matthew Davis, in progress)
- Mapping tools bias (Nuno Fonseca)
- Comparison of different quantification tools and methods (Nuno Fonseca, Natalja)
- Normalize or not-normalize: Gene Expression Atlas (Alvis Brazma, Natalja)